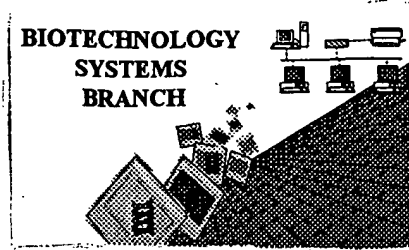


0590
0523



RAW SEQUENCE LISTING **ERROR REPORT**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/898,616
Source: API
Date Processed by STIC: 5/28/2002

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE **CHECKER**
VERSION 3.1 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND
TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom!

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addressees:

1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202
3. Hand Carry directly to:
U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202
Or
U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 01/29/2002

Raw Sequence Listing Error Summary

ERROR DETECTED

SUGGESTED CORRECTION

SERIAL NUMBER:

09/898,616

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1 Wrapped Nucleics
 Wrapped Aminos The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to 3; this will prevent "wrapping."
- 2 Invalid Line Length The rules require that a line not exceed 72 characters in length. This includes white spaces.
- 3 Misaligned Amino
 Numbering The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.
- 4 Non-ASCII The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.
- 5 Variable Length Sequence(s) contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
- 6 PatentIn 2.0
 "bug" A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) . Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.
- 7 Skipped Sequences
 (OLD RULES) Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence:
 (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
 (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
 (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
 This sequence is intentionally skipped

 Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.
- 8 Skipped Sequences
 (NEW RULES) Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence.
 <210> sequence id number
 <400> sequence id number
 000
- 9 Use of n's or Xaa's
 (NEW RULES) Use of n's and/or Xaa's have been detected in the Sequence Listing.
 Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
 In <220> to <223> section, please explain location of n or Xaa; and which residue n or Xaa represents.
- 10 Invalid <213>
 Response Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence
- 11 Use of <220> Sequence(s) missing the <220> "Feature" and associated numeric identifiers and responses.
 Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.
 (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
- 12 PatentIn 2.0
 "bug" Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
- 13 Misuse of n n can only be used to represent a single nucleotide in a nucleic acid sequence. N is not used to represent any value not specifically a nucleotide.



OIPE

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/898,616

DATE: 05/28/2002

TIME: 09:47:22

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Output Set: N:\CRF3\05282002\I898616.raw

Does Not Comply
Corrected Diskette Needed

pp 1-2, 5

of

3 <110> APPLICANT: Claragen, Inc.
 5 <120> TITLE OF INVENTION: Method for the Production of Purified rhUG for the Treatment
 6 Inflammatory and Fibrotic Conditions
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 10 <140> CURRENT APPLICATION NUMBER: 09/898,616
 11 <141> CURRENT FILING DATE: 2001-07-02
 13 <150> PRIOR APPLICATION NUMBER: 08/864,357
 14 <151> PRIOR FILING DATE: 1997-05-28
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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/898,616

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RAW SEQUENCE LISTING

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PATENT APPLICATION: US/09/898,616

TIME: 09:47:23

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RAW SEQUENCE LISTING

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338	tcatcatcag	taaccctgat	cgtgagcctc	ctctctcggt	tcatcggtat	cattaccccc	7200
340	atgaacagaa	atccccctta	cacggaggca	tcagtgaaca	aacaggaaaa	aaccgccctt	7260
342	aacatggccc	gctttatcag	aagccagaca	ttaacgcttc	tggagaaact	caacgagctg	7320
344	gacgcggatg	aacaggcaga	catctgtgaa	tcgcttcacg	accacgctga	tgagctttac	7380
346	cgcagctgcc	tcgcgcgttt	cgggtgatgac	ggtgaaaacc	tctgacacat	gcagctcccc	7440
348	gagacgggta	cagcttgtct	gtaagcggat	gccgggagca	gacaagcccc	tcaggcgcg	7500
350	tcagcgggtg	ttggcgggtg	tcggggcgca	gccatgacct	agtcacgtag	cgatagcgga	7560

RAW SEQUENCE LISTING

DATE: 05/28/2002

PATENT APPLICATION: US/09/898,616

TIME: 09:47:23

Input Set : A:\EP.txt

Output Set: N:\CRF3\05282002\I898616.raw

352 gtgtatactg gcttaactat gcggcatcag agcagattgt actgagagtg cacca 7615
355 <210> SEQ ID NO: 10
356 <211> LENGTH: 4
357 <212> TYPE: PRT
358 <213> ORGANISM: CG12
360 <400> SEQUENCE: 10
362 Ala Ala Glu Ile
363 1
365 <210> SEQ ID NO: 11
366 <211> LENGTH: 73
367 <212> TYPE: PRT
368 <213> ORGANISM: rhUG
370 <400> SEQUENCE: 11
372 Met Ala Ala Glu Ile Cys Pro Ser Phe Gln Arg Val Ile Glu Thr Leu
373 1 5 10 15
375 Leu Met Asp Thr Pro Ser Ser Tyr Glu Ala Ala Met Glu Leu Phe Ser
376 20 25 30
378 Pro Asp Gln Asp Met Arg Glu Ala Gly Ala Gln Leu Lys Lys Leu Val
379 35 40 45
381 Asp Thr Leu Pro Gln Lys Pro Arg Glu Ser Ile Ile Lys Leu Met Glu
382 50 55 60
384 Lys Ile Ala Gln Ser Ser Leu Cys Asn
385 65 70

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/898,616

DATE: 05/28/2002

TIME: 09:47:24

Input Set : A:\EP.txt

Output Set: N:\CRF3\05282002\I898616.raw